

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/539,834
Source: PCF
Date Processed by STIC: 3/14/06

ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 03/14/2006

PATENT APPLICATION: US/10/539,834

TIME: 11:06:02

Input Set : A:\10539834.txt

Output Set: N:\CRF4\03142006\J539834.raw

3 <110> APPLICANT: National Institute of Advanced Industrial Science and Technology
 4 Hisashi NARIMATSU
 5 Takashi KUDO
 6 Akira TOGAYACHI
 7 Toru HIRUMA
 9 <120> TITLE OF INVENTION: GLYCOSYLTRANSFERASE, NUCLEIC ACID ENCODING THE
 GLYCOSYLTRANSFERASE
 10 AND METHOD OF TESTING CANCERATION USING THE NUCLEIC ACID
 12 <130> FILE REFERENCE: 159-89 / YCT-902
 14 <140> CURRENT APPLICATION NUMBER: US 10/539,834
 C--> 15 <141> CURRENT FILING DATE: 2005-06-17
 17 <150> PRIOR APPLICATION NUMBER: PCT/JP03/17030
 18 <151> PRIOR FILING DATE: 2003-12-26
 20 <150> PRIOR APPLICATION NUMBER: JP 380975/2002
 21 <151> PRIOR FILING DATE: 2002-12-27
 23 <160> NUMBER OF SEQ ID NOS: 20
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 1194
 27 <212> TYPE: DNA
 28 <213> ORGANISM: Homo sapiens
 30 <400> SEQUENCE: 1
 31 atgcgctgcc ccaagtgcct tctctgcctg tcagcactgc tcacactcct gggcctcaaa 60
 32 gtgtacatcg agtggacatc cgagtcccgg ctacagcaagg cctaccccag ccctcggggc 120
 33 accccgccaa gccccacgcc agccaaccct gagcccaccc tacctgccaa cctctccacc 180
 34 cgcttgggcc agactatccc gctgcccttt gcttactgga accagcagca gtggcggtcg 240
 35 gggtccttgc ccagtgggga cagcactgaa acggggggct gccaggcttg gggggccgcc 300
 36 gccgccaccg agatccctga cttegcctcc taccccaagg acctccgccg cttcttgcctg 360
 37 tcagcagcct gccggagctt cccacagtgg ctgcctggag gtggtggcag ccaagtctcc 420
 38 agctgctcag atactgatgt cccctacctg ctgttggccg tcaagtcaga accagggcgc 480
 39 tttgcagaac gacaggccgt gagagagacg tggggcagtc cagctccagg gatccggctg 540
 40 ctcttcttgc taggggtctcc ggtgggtgag gcggggcctg acctagactc actagtggcc 600
 41 tgggagagcc gtcgctacag tgacctgctg ctctgggact tcctcgacgt cccattcaac 660
 42 cagacgctca aagacctgct gctgctggcc tggctgggcc gccactgccc caccgtgagt 720
 43 tttgtcttgc gagctcagga cgatgccttt gtacacaccc ctgccctgct ggctcacctg 780
 44 cgggccctgc cacctgcctc ggcccgaagc ctctacctgg gtgaggctct taccaggcc 840
 45 atgcctctcc ggaagccagg aggacccttc tatgtgcccg agtccttctt cgaaggtggc 900
 46 taccagcct atgcaagcgg ggggtggctac gtcattgccg ggcgcctggc accctggctg 960
 47 ctgcgggcgg cagcccgtgt ggcacccttc ccttttgagg acgtctacac tggcctttgc 1020
 48 atccgagccc tgggcctggt gccccaggcc caccaggtct tcctcacagc ctggccagca 1080
 49 gaccgcactg cggaccactg tgctttccgc aacctgctgc tggtaaggcc cctggggccc 1140
 50 caggccagca ttcggctctg gaaacaactg caagacccaa ggctccagtg ctga 1194
 52 <210> SEQ ID NO: 2
 53 <211> LENGTH: 397
 54 <212> TYPE: PRT

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55 <213> ORGANISM: Homo sapiens

57 <400> SEQUENCE: 2

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58 Met Arg Cys Pro Lys Cys Leu Leu Cys Leu Ser Ala Leu Leu Thr Leu
59 1          5          10          15
60 Leu Gly Leu Lys Val Tyr Ile Glu Trp Thr Ser Glu Ser Arg Leu Ser
61          20          25          30
62 Lys Ala Tyr Pro Ser Pro Arg Gly Thr Pro Pro Ser Pro Thr Pro Ala
63          35          40          45
64 Asn Pro Glu Pro Thr Leu Pro Ala Asn Leu Ser Thr Arg Leu Gly Gln
65          50          55          60
66 Thr Ile Pro Leu Pro Phe Ala Tyr Trp Asn Gln Gln Trp Arg Leu
67 65          70          75          80
68 Gly Ser Leu Pro Ser Gly Asp Ser Thr Glu Thr Gly Gly Cys Gln Ala
69          85          90          95
70 Trp Gly Ala Ala Ala Ala Thr Glu Ile Pro Asp Phe Ala Ser Tyr Pro
71          100         105         110
72 Lys Asp Leu Arg Arg Phe Leu Leu Ser Ala Ala Cys Arg Ser Phe Pro
73          115         120         125
74 Gln Trp Leu Pro Gly Gly Gly Gly Ser Gln Val Ser Ser Cys Ser Asp
75          130         135         140
76 Thr Asp Val Pro Tyr Leu Leu Leu Ala Val Lys Ser Glu Pro Gly Arg
77 145          150          155          160
78 Phe Ala Glu Arg Gln Ala Val Arg Glu Thr Trp Gly Ser Pro Ala Pro
79          165          170          175
80 Gly Ile Arg Leu Leu Phe Leu Leu Gly Ser Pro Val Gly Glu Ala Gly
81          180          185          190
82 Pro Asp Leu Asp Ser Leu Val Ala Trp Glu Ser Arg Arg Tyr Ser Asp
83          195         200         205
84 Leu Leu Leu Trp Asp Phe Leu Asp Val Pro Phe Asn Gln Thr Leu Lys
85          210         215         220
86 Asp Leu Leu Leu Leu Ala Trp Leu Gly Arg His Cys Pro Thr Val Ser
87 225          230         235         240
88 Phe Val Leu Arg Ala Gln Asp Asp Ala Phe Val His Thr Pro Ala Leu
89          245         250         255
90 Leu Ala His Leu Arg Ala Leu Pro Pro Ala Ser Ala Arg Ser Leu Tyr
91          260         265         270
92 Leu Gly Glu Val Phe Thr Gln Ala Met Pro Leu Arg Lys Pro Gly Gly
93          275         280         285
94 Pro Phe Tyr Val Pro Glu Ser Phe Phe Glu Gly Gly Tyr Pro Ala Tyr
95          290         295         300
96 Ala Ser Gly Gly Gly Tyr Val Ile Ala Gly Arg Leu Ala Pro Trp Leu
97 305          310         315         320
98 Leu Arg Ala Ala Ala Arg Val Ala Pro Phe Pro Phe Glu Asp Val Tyr
99          325         330         335
100 Thr Gly Leu Cys Ile Arg Ala Leu Gly Leu Val Pro Gln Ala His Pro
101          340         345         350
102 Gly Phe Leu Thr Ala Trp Pro Ala Asp Arg Thr Ala Asp His Cys Ala
103          355         360         365
104 Phe Arg Asn Leu Leu Leu Val Arg Pro Leu Gly Pro Gln Ala Ser Ile

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105      370      375      380
106 Arg Leu Trp Lys Gln Leu Gln Asp Pro Arg Leu Gln Cys
107 385      390      395      397
110 <210> SEQ ID NO: 3
111 <211> LENGTH: 31
112 <212> TYPE: DNA
113 <213> ORGANISM: Artificial Sequence
115 <220> FEATURE:
116 <223> OTHER INFORMATION: Description of Artificial Sequence: 5' primer for PCR
118 <400> SEQUENCE: 3
119 ctcaagctta tgcgctgccc caagtgcctt c 31
121 <210> SEQ ID NO: 4
122 <211> LENGTH: 31
123 <212> TYPE: DNA
124 <213> ORGANISM: Artificial Sequence
126 <220> FEATURE:
127 <223> OTHER INFORMATION: Description of Artificial Sequence: 3' primer for PCR
129 <400> SEQUENCE: 4
130 ctgaattct cagcactgga gccttggggtc t 31
132 <210> SEQ ID NO: 5
133 <211> LENGTH: 20
134 <212> TYPE: DNA
135 <213> ORGANISM: Artificial Sequence
137 <220> FEATURE:
138 <223> OTHER INFORMATION: Description of Artificial Sequence: 5' primer for RT-PCR
140 <400> SEQUENCE: 5
141 gctgttggcc gtcaagtcag 20
143 <210> SEQ ID NO: 6
144 <211> LENGTH: 18
145 <212> TYPE: DNA
146 <213> ORGANISM: Artificial Sequence
148 <220> FEATURE:
149 <223> OTHER INFORMATION: Description of Artificial Sequence: 3' primer for RT-PCR
151 <400> SEQUENCE: 6
152 caggaagagc agccggat 18
154 <210> SEQ ID NO: 7
155 <211> LENGTH: 18
156 <212> TYPE: DNA
157 <213> ORGANISM: Artificial Sequence
159 <220> FEATURE:
160 <223> OTHER INFORMATION: Description of Artificial Sequence: probe for RT-PCR
162 <400> SEQUENCE: 7
163 cagaacgaca ggccgtga 18
165 <210> SEQ ID NO: 8
166 <211> LENGTH: 29
167 <212> TYPE: DNA
168 <213> ORGANISM: Artificial Sequence
170 <220> FEATURE:
171 <223> OTHER INFORMATION: Description of Artificial Sequence: 5' primer for PCR

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Input Set : A:\10539834.txt

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173 <400> SEQUENCE: 8
174 gccaaagctta catccgagtc ccggctcag                29
176 <210> SEQ ID NO: 9
177 <211> LENGTH: 29
178 <212> TYPE: DNA
179 <213> ORGANISM: Artificial Sequence
181 <220> FEATURE:
182 <223> OTHER INFORMATION: Description of Artificial Sequence: 5' primer for PCR
184 <400> SEQUENCE: 9
185 gccaaagctta aggcctaccc cagccctcg                29
187 <210> SEQ ID NO: 10
188 <211> LENGTH: 28
189 <212> TYPE: DNA
190 <213> ORGANISM: Artificial Sequence
192 <220> FEATURE:
193 <223> OTHER INFORMATION: Description of Artificial Sequence: 3' primer for PCR
195 <400> SEQUENCE: 10
196 cggaattctc agcactggag ccttgggt                28
198 <210> SEQ ID NO: 11
199 <211> LENGTH: 55
200 <212> TYPE: DNA
201 <213> ORGANISM: Artificial Sequence
203 <220> FEATURE:
204 <223> OTHER INFORMATION: Description of Artificial Sequence: 5' primer for PCR
206 <400> SEQUENCE: 11
207 ggggacaagt ttgtacaaaa aagcaggctt ccccagccct cggggcaccc cgcca                55
209 <210> SEQ ID NO: 12
210 <211> LENGTH: 54
211 <212> TYPE: DNA
212 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: Description of Artificial Sequence: 3' primer for PCR
217 <400> SEQUENCE: 12
218 ggggaccact ttgtacaaga aagctgggtc tcagcactgg agccttgggt cttg                54
220 <210> SEQ ID NO: 13
221 <211> LENGTH: 29
222 <212> TYPE: DNA
223 <213> ORGANISM: Artificial Sequence
225 <220> FEATURE:
226 <223> OTHER INFORMATION: Description of Artificial Sequence: 5' primer for PCR
228 <400> SEQUENCE: 13
229 gccaaagctta catccgagtc ccggctcag                29
231 <210> SEQ ID NO: 14
232 <211> LENGTH: 29
233 <212> TYPE: DNA
234 <213> ORGANISM: Artificial Sequence
236 <220> FEATURE:
237 <223> OTHER INFORMATION: Description of Artificial Sequence: 5' primer for PCR
239 <400> SEQUENCE: 14

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RAW SEQUENCE LISTING

DATE: 03/14/2006

PATENT APPLICATION: US/10/539,834

TIME: 11:06:03

Input Set : A:\10539834.txt

Output Set: N:\CRF4\03142006\J539834.raw

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240 gccaaagctta aggcctaccc cagccctcg                               29
242 <210> SEQ ID NO: 15
243 <211> LENGTH: 28
244 <212> TYPE: DNA
245 <213> ORGANISM: Artificial Sequence
247 <220> FEATURE:
248 <223> OTHER INFORMATION: Description of Artificial Sequence: 3' primer for PCR
250 <400> SEQUENCE: 15
251 cggaattctc agcactggag ccttgggt                               28
254 <210> SEQ ID NO: 16
255 <211> LENGTH: 372
256 <212> TYPE: PRT
257 <213> ORGANISM: Homo sapiens
259 <400> SEQUENCE: 16
260 Thr Ser Glu Ser Arg Leu Ser Lys Ala Tyr Pro Ser Pro Arg Gly Thr
261 1                               5                               10                               15
262 Pro Pro Ser Pro Thr Pro Ala Asn Pro Glu Pro Thr Leu Pro Ala Asn
263                               20                               25                               30
264 Leu Ser Thr Arg Leu Gly Gln Thr Ile Pro Leu Pro Phe Ala Tyr Trp
265                               35                               40                               45
266 Asn Gln Gln Gln Trp Arg Leu Gly Ser Leu Pro Ser Gly Asp Ser Thr
267                               50                               55                               60
268 Glu Thr Gly Gly Cys Gln Ala Trp Gly Ala Ala Ala Thr Glu Ile
269 65                               70                               75                               80
270 Pro Asp Phe Ala Ser Tyr Pro Lys Asp Leu Arg Arg Phe Leu Leu Ser
271                               85                               90                               95
272 Ala Ala Cys Arg Ser Phe Pro Gln Trp Leu Pro Gly Gly Gly Ser
273                               100                              105                              110
274 Gln Val Ser Ser Cys Ser Asp Thr Asp Val Pro Tyr Leu Leu Ala
275                               115                              120                              125
276 Val Lys Ser Glu Pro Gly Arg Phe Ala Glu Arg Gln Ala Val Arg Glu
277                               130                              135                              140
278 Thr Trp Gly Ser Pro Ala Pro Gly Ile Arg Leu Leu Phe Leu Leu Gly
279 145                              150                              155                              160
280 Ser Pro Val Gly Glu Ala Gly Pro Asp Leu Asp Ser Leu Val Ala Trp
281                               165                              170                              175
282 Glu Ser Arg Arg Tyr Ser Asp Leu Leu Leu Trp Asp Phe Leu Asp Val
283                               180                              185                              190
284 Pro Phe Asn Gln Thr Leu Lys Asp Leu Leu Leu Leu Ala Trp Leu Gly
285                               195                              200                              205
286 Arg His Cys Pro Thr Val Ser Phe Val Leu Arg Ala Gln Asp Asp Ala
287                               210                              215                              220
288 Phe Val His Thr Pro Ala Leu Leu Ala His Leu Arg Ala Leu Pro Pro
289 225                              230                              235                              240
290 Ala Ser Ala Arg Ser Leu Tyr Leu Gly Glu Val Phe Thr Gln Ala Met
291                               245                              250                              255
292 Pro Leu Arg Lys Pro Gly Gly Pro Phe Tyr Val Pro Glu Ser Phe Phe
293                               260                              265                              270
294 Glu Gly Gly Tyr Pro Ala Tyr Ala Ser Gly Gly Gly Tyr Val Ile Ala

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VERIFICATION SUMMARY

DATE: 03/14/2006

PATENT APPLICATION: US/10/539,834

TIME: 11:06:04

Input Set : A:\10539834.txt

Output Set: N:\CRF4\03142006\J539834.raw

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date